```
                        10         20         30         40         50
DSFBP314.AMI      1 MSNRPIYLDY SATTPVDPSV VEKMIPWLYE SFGNPASRSH AFGWEAEDAV   50
DSFBP536.AMI      1 MSNRPIYLDY SATTPVDPSV VEKMIPWLYE SFGNPASRSH AFGWEAEDAV   50
                        60         70         80         90        100
DSFBP314.AMI     51 EKAREEVAKL VNADPREIVW TSGATESDNL AIKGAANFYA ERGKHIITVK  100
DSFBP536.AMI     51 EKAREEVAKL VNADPREIVW TSGATESDNL AIKGAANFYA ERGKHIITVK  100
                       110        120        130        140        150
DSFBP314.AMI    101 TEHKAVLDTC RELERQGFEV TYLDVQDDGL LSLDAFKAAL RPDTILVSVM  150
DSFBP536.AMI    101 TEHKAVLDTC RELERQGFEV TYLDVQDDGL LSLDAFKAAL RPDTILVSVM  150
                       160        170        180        190        200
DSFBP314.AMI    151 MVNNEIGVIQ DIAALGEICR EKGIIFHVDA AQATGKVEID LQKLKVDLMS  200
DSFBP536.AMI    151 MVNNEIGVIQ DIAALGEICR EKGIIFHVDA AQATGKVEID LQKLKVDLMS  200
                       210        220        230        240        250
DSFBP314.AMI    201 FSAHKTYGPK GIGALYVRRK PRVRIEAQMH GGGHERGFRS GTLATHQIVG  250
DSFBP536.AMI    201 FSAHKTYGPK GIGALYVRRK PRVRIEAQMH GGGHERGFRS GTLATHQIVG  250
                       260        270        280        290        300
DSFBP314.AMI    251 MGEAFRLARE EMGTENERVR MLRDRLLAGL TQIEEVYVNG SMEHRVPHNL  300
DSFBP536.AMI    251 MGEAFRLARE EMGTENERVR MLRDRLLAGL TQIEEVYVNG SMEHRVPHNL  300
                       310        320        330        340        350
DSFBP314.AMI    301 NISFNYVEGE SLIMAIKELA VSSGSACTSA SLEPSYVLRA LGRNDELAHS  350
DSFBP536.AMI    301 NISFNYVEGE SLIMAIKELA VSSGSACTSA SLEPSYVLRA LGRNDELAHS  350
                       360        370        380        390        400
DSFBP314.AMI    351 SIRFTLGRFT TEQEIDFTIE LIKSRVGKLR DMSPLWEMAQ EGIDLNSVQW  400
DSFBP536.AMI    351 SIRFTLGRFT TEQEIDFTIE LIKSRVGKLR DMSPLWEMAQ EGIDLNSVQW  400
                       410        420        430        440        450
DSFBP314.AMI    401 AAH*..... .......... .......... .......... ..........  450
DSFBP536.AMI    401 AAH*..... .......... .......... .......... ..........  450
                        10         20         30         40         50
DSF314.DNA        1 ATGAGCAATC GCCCCATCTA CCTGGACTAC TCGGCTACCA CGCCGGTCGA   50
DSF536F1.DNA      1 ATGAGCAATC GCCCCATCTA CCTGGACTAC TCGGCTACCA CGCCGGTCGA   50
DSF536R1.DNA      1 ---------- ---------- ---------- ---------- ----------   50
DSF53611.DNA      1 ---------- ---------- ---------- ---------- ----------   50
DSF53612.DNA      1 ---------- ---------- ---------- ---------- ----------   50
                        60         70         80         90        100
DSF314.DNA       51 CCCGAGCGTG GTCGAGAAAA TGATTCCCTG GTTGTACGAG AGTTTCGGCA  100
DSF536F1.DNA     51 CCCGAGCGTG GTCGAGAAAA TGATTCCCTG GTTGTACGAG AGTTTCGGCA  100
DSF536R1.DNA     51 ---------- ---------- ---------- ---------- ----------  100
DSF53611.DNA     51 ---------- ---------- ---------- ---------- ----------  100
DSF53612.DNA     51 ---------- ---------- ---------- ---------- ----------  100
                       110        120        130        140        150
DSF314.DNA      101 ATCCGGCCTC GCGCAGCCAC GCCTTTGGCT GGGAAGCCGA GGACGCGGTC  150
DSF536F1.DNA    101 ATCCGGCCTC GCGCAGCCAC GCCTTTGGCT GGGAAGCCGA GGACGCGGTC  150
DSF536R1.DNA    101 ---------- ---------- ---------- ---------- ----------  150
DSF53611.DNA    101 ---------- ---------- ---------- ---------- ----------  150
DSF53612.DNA    101 ---------- ---------- ---------- ---------- ----------  150
                       160        170        180        190        200
DSF314.DNA      151 GAGAAGGCCC GCGAGGAAGT TGCCAAGCTG GTCAACGCCG ATCCGCGCGA  200
DSF536F1.DNA    151 GAGAAGGCCC GCGAGGAAGT TGCCAAGCTG GTCAACGCCG ATCCGCGCGA  200
DSF536R1.DNA    151 ---------- ---------- ---------- ---------- ----------  200
DSF53611.DNA    151 ---------- ---------- ---------- ---------- ----------  200
DSF53612.DNA    151 ---------- ---------- ---------- ---------- ----------  200
                       210        220        230        240        250
DSF314.DNA      201 GATCGTCTGG ACTTCCGGCG CTACCGAGTC GGACAACCTG GCCATCAAGG  250
DSF536F1.DNA    201 GATCGTCTGG ACTTCCGGCG CTACCGAGTC GGACAACCTG GCCATCAAGG  250
DSF536R1.DNA    201 ---------- ---------- ---------- ---------- ----------  250
DSF53611.DNA    201 ---------- ---------- ---------- ---------- ----------  250
DSF53612.DNA    201 ---------- ---------- ---------- ---------- ----------  250
                       260        270        280        290        300
DSF314.DNA      251 GCGCGGCGAA TTTCTACGCC GAGCGCGGCA AGCACATCAT CACCGTCAAG  300
DSF536F1.DNA    251 GCGCGGCGAA TTTCTACGCC GAGCGCGGCA AGCACATCAT CACCGTCAAG  300
DSF536R1.DNA    251 ---------- ---------- ---------- ---------- ----------  300
DSF53611.DNA    251 ---------- ---------- ---------- ---------- ----------  300
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Figure 7A

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DSF53612.DNA    251 ---------- ---------- ---------- ---------- ----------  300
                          310        320        330        340        350
DSF314.DNA      301 ACCGAACACA AGGCGGTGCT GGATACCTGT CGGGAGCTCG AACGCCAGGG  350
DSF536F1.DNA    301 ACCGAACACA AGGCGGTGCT GGATACCTGT CGGGAGCTCG AACGCCAGGG  350
DSF536R1.DNA    301 ---------- ---------- ---------- ---------- ----------  350
DSF53611.DNA    301 ---------- ---------- ---------- ---------- ----------  350
DSF53612.DNA    301 ---------- ---------- ---------- ---------- ----------  350
                          360        370        380        390        400
DSF314.DNA      351 CTTTGAAGTG ACCTACCTGG ATGTCCAGGA CGATGGTCTG CTCAGCCTCG  400
DSF536F1.DNA    351 CTTTGAAGTG ACCTACCTGG ATGTCCAGGA CGATGGTCTG CTCAGCCTCG  400
DSF536R1.DNA    351 ---------- ---------- ---------- ---------- ----------  400
DSF53611.DNA    351 ---------- ---------- ---------- ---------- ----------  400
DSF53612.DNA    351 ---------- ---------- ---------- ---------- ----------  400
                          410        420        430        440        450
DSF314.DNA      401 ATGCGTTCAA GGCTGCGCTG CGCCCGGATA CCATCCTGGT GTCGGTGATG  450
DSF536F1.DNA    401 ATGCGTTCAA GGCTGCGCTG CGCCCGGATA CCATCCTGGT GTCGGTGATG  450
DSF536R1.DNA    401 ---------- ---------- ---------- ---------- ----------  450
DSF53611.DNA    401 ---------- ---------- ---------- ---------- ----------  450
DSF53612.DNA    401 ---------- ---------- ---------- ----CCTGGT GTCGGTGATG  450
                          460        470        480        490        500
DSF314.DNA      451 ATGGTCAACA ACGAGATCGG CGTCATCCAG GACATCGCCG CGCTGGGCGA  500
DSF536F1.DNA    451 ATGGTCAACA ACGAGATCGG CGTCATCCAG GACATCGCCG CGCTGGGCGA  500
DSF536R1.DNA    451 ---------- ---------- ---------- ---------- ----------  500
DSF53611.DNA    451 ---------- ---------- ---------- ---------- ----------  500
DSF53612.DNA    451 ATGGTCAACA ACGAGATCGG CGTCATCCAG GACATCGCCG CGCTGGGCGA  500
                          510        520        530        540        550
DSF314.DNA      501 GATCTGCCGC GAGAAGGGCA TCATCTTCCA CGTGGACGCG GCCCAGGCCA  550
DSF536F1.DNA    501 GATCTGCCGC GAGAAGGGCA -CATCTTCCA CGTGGACGCG GCC-AAGCCA  550
DSF536R1.DNA    501 ---------- ---------- ---------- ---------- ----------  550
DSF53611.DNA    501 ---------- ---------- ---------- ---------- ---------C  550
DSF53612.DNA    501 GATCTGCCGC GAGAAGGGCA TCATCTTCCA CGTGGACGCG GCCCAGGCCA  550
                          560        570        580        590        600
DSF314.DNA      551 CCGGCAAGGT CGAGATCGAC CTGCAGAAGC TGAAGGTGGA CCTGATGTCG  600
DSF536F1.DNA    551 ACGGCAAGGT CGAGATC--- ---------- ---------- ----------  600
DSF536R1.DNA    551 ---------- ---------- ---------- ---------- ----------  600
DSF53611.DNA    551 ---------- -----TCGAC CTGCAGAAGC TGAAGGTGGA CCTGATGTCG  600
DSF53612.DNA    551 CCGGCAAGGT CGAGATCGAC CTGCAGAAGC TGAAGGTGGA CCTGATGTCG  600
                          610        620        630        640        650
DSF314.DNA      601 TTCTCGGCGC ACAAGACGTA CGGCCCCAAG GGCATCGGCG CGCTGTATGT  650
DSF536F1.DNA    601 ---------- ---------- ---------- ---------- ----------  650
DSF536R1.DNA    601 ---------- ---------- ---------- ---------- ----------  650
DSF53611.DNA    601 TTCTCGGCGC ACAAGACGTA CGGCCCCAAG GGCATCGGCG CGCTGTATGT  650
DSF53612.DNA    601 TTCTCGGCGC ACAAGACGTA CGGCCCCAAG GGCATCGGCG CGCTGTATGT  650
                          660        670        680        690        700
DSF314.DNA      651 GCGGCGCAAG CCGCGCGTGC GCATCGAGGC GCAGATGCAC GGCGGCGGCC  700
DSF536F1.DNA    651 ---------- ---------- ---------- ---------- ----------  700
DSF536R1.DNA    651 --GGCGCAAG CCGCGCGTGN GNATCGAGGC GCAGATGCAC GGCGGCGGCC  700
DSF53611.DNA    651 GCGGCGCAAG CCGCGCGTGC GCATCGAGGC GCAGATGCAC GGCGGCGGCC  700
DSF53612.DNA    651 GCGGCGCAAG CCGCGCGTGC GCATCGAGGC NTAGATGCAC GGCGGCGGCC  700
                          710        720        730        740        750
DSF314.DNA      701 ACGAACGGGG CTTCCGGTCG GGCACGCTGG CCACGCACCA GATCGTCGGC  750
DSF536F1.DNA    701 ---------- ---------- ---------- ---------- ----------  750
DSF536R1.DNA    701 ACGAACGGGG CTTCCGGTCG GGCACGNTGG CCACGCACCA GATCGTCGGC  750
DSF53611.DNA    701 ACGAACGGGG CTTCCGGTCG GGCACGCTGG CCACGCACCA GATCGTCGGC  750
DSF53612.DNA    701 ACGAACG--- ---------- ---------- ---------- ----------  750
                          760        770        780        790        800
DSF314.DNA      751 ATGGGCGAGG CGTTCCGCCT GGCGCGCGAG GAAATGGGCA CCGAGAACGA  800
DSF536F1.DNA    751 ---------- ---------- ---------- ---------- ----------  800
DSF536R1.DNA    751 ATGGGCGAGG CGTTCCGCCT GGCGCGCGAG GAAATGGGCA CCGAGAACGA  800
DSF53611.DNA    751 ATGGGCGAGG CGTTCCGCCT GGCGCGCGAG GAAATGGGCA CCGAGAACGA  800
DSF53612.DNA    751 ---------- ---------- ---------- ---------- ----------  800
                          810        820        830        840        850
DSF314.DNA      801 GCGCGTGCGC ATGCTGCGCG ACCGCCTGCT GGCCGGCCTG ACGCAGATCG  850
DSF536F1.DNA    801 ---------- ---------- ---------- ---------- ----------  850
DSF536R1.DNA    801 GCGCGTGCGC ATGCTGCGCG ACCGCCTGCT GGCCGGCCTG ACGCAGATCG  850
DSF53611.DNA    801 GCGCGTGCGC ATGCTGCGCG ACCGCCTGCT GGCCGGCCTG ACGCAGATCG  850
DSF53612.DNA    801 ---------- ---------- ---------- ---------- ----------  850
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Figure 7B

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                      860        870        880        890        900
DSF314.DNA      851 AGGAAGTGTA TGTGAACGGC AGCATGGAGC ACCGCGTGCC GCACAACCTG 900
DSF536F1.DNA    851 ---------- ---------- ---------- ---------- ---------- 900
DSF536R1.DNA    851 AGGAAGTGTA TGTGAACGGC AGCATGGAGC ACCGCGTGCC GCACAACCTG 900
DSF53611.DNA    851 AGGAAGTGTA TGTGAACGGC AGCATGGAGC ACCGCGTGCC GCACAACCTG 900
DSF53612.DNA    851 ---------- ---------- ---------- ---------- ---------- 900
                      910        920        930        940        950
DSF314.DNA      901 AACATCAGCT TCAACTATGT CGAGGGCGAG TCTCTGATCA TGGCGATCAA 950
DSF536F1.DNA    901 ---------- ---------- ---------- ---------- ---------- 950
DSF536R1.DNA    901 AACATCAGCT TCAACTATGT CGAGGGCGAG TCTCTGATCA TGGCGATCAA 950
DSF53611.DNA    901 AACATCAGCT TCAACTATGT CGAGGGCGAG TCTCTGATCA TGGCGATCAA 950
DSF53612.DNA    901 ---------- ---------- ---------- ---------- ---------- 950
                      960        970        980        990       1000
DSF314.DNA      951 GGAGCTGGCC GTTTCCAGCG GTTCGGCCTG CACGTCGGCC AGCCTGGAGC 1000
DSF536F1.DNA    951 ---------- ---------- ---------- ---------- ---------- 1000
DSF536R1.DNA    951 GGAGCTGGCC GTTTCCAGCG GTTCGGCCTG CACGTCGGCN AGCCTGGAGC 1000
DSF53611.DNA    951 GGAGCTGGCC GTTTCCAGCG GTTCGGCCTG CACGTCGGC- ---------- 1000
DSF53612.DNA    951 ---------- ---------- ---------- ---------- ---------- 1000
                     1010       1020       1030       1040       1050
DSF314.DNA     1001 CGTCCTATGT GCTGCGCGCG CTGGGCCGCA ACGACGAGCT GGCGCACAGC 1050
DSF536F1.DNA   1001 ---------- ---------- ---------- ---------- ---------- 1050
DSF536R1.DNA   1001 CGTCCTATGT GCTGCGCGCG CTGGGCCGCA ACGACGAGCT GGCGCACAGC 1050
DSF53611.DNA   1001 ---------- ---------- ---------- ---------- ---------- 1050
DSF53612.DNA   1001 ---------- ---------- ---------- ---------- ---------- 1050
                     1060       1070       1080       1090       1100
DSF314.DNA     1051 TCCATCCGCT TTACCCTGGG CCGCTTCACG ACCGAACAGG AAATCGACTT 1100
DSF536F1.DNA   1051 ---------- ---------- ---------- ---------- ---------- 1100
DSF536R1.DNA   1051 TCCATCCGCT TTACCCTGGG CCGCTTCACG ACCGAACAGG AAATCGACTT 1100
DSF53611.DNA   1051 ---------- ---------- ---------- ---------- ---------- 1100
DSF53612.DNA   1051 ---------- ---------- ---------- ---------- ---------- 1100
                     1110       1120       1130       1140       1150
DSF314.DNA     1101 CACGATCGAA CTGATCAAGA GTCGTGTCGG CAAGCTGCGC GATATGTCGC 1150
DSF536F1.DNA   1101 ---------- ---------- ---------- ---------- ---------- 1150
DSF536R1.DNA   1101 CACGATCGAA CTGATCAAGA GTCGTGTCGG CAAGCTGCGC GATATGTCGC 1150
DSF53611.DNA   1101 ---------- ---------- ---------- ---------- ---------- 1150
DSF53612.DNA   1101 ---------- ---------- ---------- ---------- ---------- 1150
                     1160       1170       1180       1190       1200
DSF314.DNA     1151 CGTTGTGGGA AATGGCCCAG GAAGGCATTG ATCTGAATTC CGTGCAGTGG 1200
DSF536F1.DNA   1151 ---------- ---------- ---------- ---------- ---------- 1200
DSF536R1.DNA   1151 CGTTGTGGGA AATGGCCCAG GAAGGCATTG ATCTGAATTC CGTGCAGTGG 1200
DSF53611.DNA   1151 ---------- ---------- ---------- ---------- ---------- 1200
DSF53612.DNA   1151 ---------- ---------- ---------- ---------- ---------- 1200
                     1210       1220       1230       1240       1250
DSF314.DNA     1201 GCCGCGCACT GA........ .......... .......... .......... 1250
DSF536F1.DNA   1201 ---------- -......... .......... .......... .......... 1250
DSF536R1.DNA   1201 GCCGCGCACT GA........ .......... .......... .......... 1250
DSF53611.DNA   1201 ---------- -......... .......... .......... .......... 1250
DSF53612.DNA   1201 ---------- -......... .......... .......... .......... 1250
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Figure 7C